

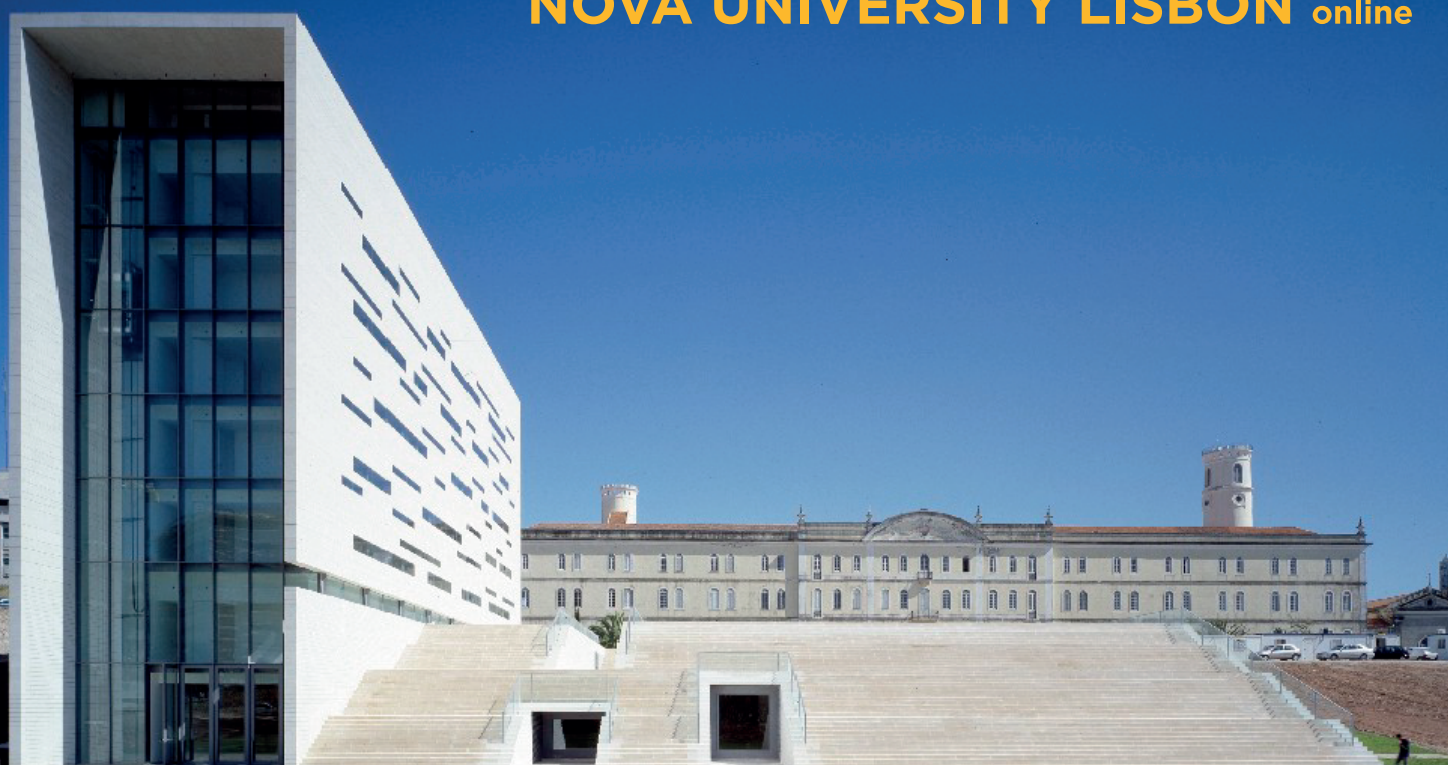
MICROBIOTECH 21

Webconference

CONGRESS OF MICROBIOLOGY AND BIOTECHNOLOGY 2021

November 23rd - 26th | 2021

NOVA UNIVERSITY LISBON online



Abstracts Book

www.microbiotec21.organideia.pt



Organizing Committee	16
Scientific Committee	16
PCO Secretariat	18
Plenary Presentations	19
Topic #1: Microbiotech in Health & Disease	19
SARS-CoV-2 variants - the steering wheel of public health and governments	19
3D culture strategies to address cell microenvironment remodeling in disease and therapeutic response	20
Topic #2: Microbiotech for a Sustainable Future	21
Potentials of synthetic biology for fine and speciality chemical production	21
Plenary 4 – Sunil Chandran (Amyris Biotechnologies)	22
Topic #3: Food and Environmental Microbiotech	23
Antimicrobial resistance in the ecosystem: some recent advances	23
Topic #4: Impact of Microbiotech in Society	24
Startup Lessons from a Bio-Entrepreneur Turned Investor	24
Topic #5: Cool tools in Microbiotech	25
Electrochemical detection of single nucleotide polymorphism associated with rifampicin resistance in Mycobacterium tuberculosis using solid-phase primer elongation with ferrocene-linked redox-labelled nucleotides	25
Antibiotic translocation across bacterial membranes and bacterial susceptibility, concepts & methods	27
Keynotes	29
1.1 Zoonoses, emerging and re-emerging diseases: virology, bacteriology, mycology and parasitology	29
SARS-CoV2 a threat across the human-animal-environment dimensions: how the prediction, prevention, detection and response to emerging infectious diseases needs a new technological boost	29
1.2. Microbiome-in-health-and-disease-microbial-diversity-and-host-microbes-interactions	31
Pathogens and the microbiome: antibiotics at the crossroad	31
1.3 Microbiology & Biotechnology in COVID-19: new approaches for new diseases	32
Signatures in SARS-CoV-2 spike protein conferring escape to neutralizing antibodies	32
1.4 & 1.6 Biopharmaceuticals & Vaccines: design, analytical methods and bioprocessing opportunities	33
Two-Component Nanoparticle Vaccine Displaying Glycosylated Spike S1 Domain Induces Neutralizing Antibody Response against SARS-CoV-2 Variants	33
RNA as biopharmaceuticals – the challenges behind bioprocessing	34
1.5 Antimicrobial Resistance in One Health: the silent pandemics	35
Antimicrobial resistance in One Health: the silent pandemic on human health	35
Antimicrobial resistance in One Health: the silent pandemic on the animal and environmental settings	36
1.7 Molecular microbiology in the NGS era: early detection and identification of pathogens and its dispersion and evolution	37
How next-generation sequencing (NGS) is helping to preview, prevent, and manage public health threats	37
2.1 Biotechnology and Sustainability I	39
Biotechnology in circular economy & bioeconomy: Bioengineering and microbiology for sustainable development: the case of bioethanol	39
2.2 Biotechnology and Sustainability II	41
Hybrid High Throughput Process Development for Biologics Production	41
2.3 Biotechnology and Sustainability III	43

New phototrophic factories for nutrient recovery and polyhydroxyalkanoates production	43
3.1 Microbial physiology and biochemistry: microbial activity in the context of environmental changes	45
A novel capsular polysaccharide from <i>Shewanella vesiculosa</i> HM13 with adhesive properties	45
3.2 Biotechnology and environmental microbiology: from psychrophiles to extremophiles	46
Diversity, secondary metabolites and biotechnological potential of marine-derived actinomycetes from the coasts of Continental Portugal and the Macaronesia archipelagos	46
3.3 Microbiological collections: resources for a sustainable present and future	48
Creating novel opportunities for using microbial resources	48
3.4 Soil health, food security and food safety	50
Fresh produce bioprotection and biopreservation	50
3.5 Biotech4food – nutrigenomics, personalized nutrition and food production	51
Environment, gene and personalized nutrition	51
4.1 Teaching and Communicating in Science	52
The International Microbiology Literacy Initiative	52
4.2 COST sessions rooms	54
COST (European Cooperation in Science and Technology): Funding networking opportunities	54
5.1 Genome Editing for therapy, biotechnological & food use	55
Pedro Fevereiro - TBA	55
5.2 Synthetic & Systems biology: experimental, theoretical and computational approaches for the analysis of micro and biological systems	56
Synthetic biology approaches to engineer polyphenols microbial cell factories	56
5.3 New approaches for disease control and treatment	57
Pulmonary Phage Therapy: from proof of concept to pre-clinical assessment	57
Targeting viral ribonucleases from SARS-CoV-2 to combat COVID-19	58
5.4 Digital Data, Big Data & AI in Microbiotech	59
<i>In data speramus</i> – How do deliver against the believe in the power of machine learning in biotechnology	59
Oral Presentations	60
1.1 Zoonoses, emerging and re-emerging diseases: virology, bacteriology, mycology and parasitology	60
O.167. Live cell biosensors for detection and quantification of viral pathogens	60
O.257. The role of Toll Like Receptors on the innate immune response against <i>Mycobacterium ulcerans infection</i>	61
O.361. A new high-performance method to detect, identify, and sort metabolically active cells of environmental <i>Mycobacterium bovis</i>	63
1.2 Microbiome in Health & Disease: microbial diversity and host-microbes interactions	65
O.1. Multiple acyl-CoA dehydrogenase deficiency kills <i>Mycobacterium tuberculosis</i> in vitro and during infection	65
O.24. Pneumococcal colonization, smoking status and contact with children influence the upper respiratory tract microbiota of healthy adults	66
O.67. A new <i>Legionella pneumophila</i> effector displaying strain-specific nucleotropism	68
O.110. Identification, phenotypic characterization, and selection of gilthead seabream (<i>Sparus aurata</i>) associated bacteria for application as putative probiotics in fish larviculture	69
O.139. Efficient encapsulation of the newly synthesized cell wall protects <i>Streptococcus pneumoniae</i> from peptidoglycan hydrolases and host defenses	70
O.157. Virulence gene repression promotes <i>Listeria monocytogenes</i> systemic infection	71
O.189. <i>Burkholderia cepacia</i> complex silencing proteins as mediators of horizontally transferred genes expression during infection of host cells	72
O.280. Characterization of growth and quorum sensing responses in <i>Barnesiella</i> and <i>Muribaculum</i> gut microbiota members	74

1.3 Microbiology & Biotechnology in COVID-19: new approaches for new diseases	76
O.6. SARS-CoV-2 membrane protein: from genomic data to structural new insights	76
O.272. Wastewater monitoring of SARS-CoV-2 RNA in Porto, Portugal: temporal variations during the COVID-19 pandemic	78
O.406. SYBR Green-based RT-PCR tests for rapid, sensitive and inexpensive detection of SARS-CoV-2	80
1.4 & 1.6 - Biopharmaceuticals & Vaccines: design, analytical methods and bioprocessing opportunities	81
O.49. Development of recombinant vaccines against <i>Vibrio</i> spp. using an outer membrane channel as target	81
O.92. Cyanobacterial extracellular vesicles as novel molecular-display platforms targeting fish-related applications	83
O.296. A cyanobacterial extracellular carbohydrate polymer with strong antitumor activity	85
O.388. Biocompatible excipients to improve the stability of immunoglobulin Y (IgY) antibodies	87
O.412. Cell line engineering for stable expression of cytotoxic envelope glycoproteins to enable continuous lentiviral vector production	89
O.439. Minicircle-derived RNA interference as a novel gene-based anti-angiogenic therapy	90
1.5 Antimicrobial Resistance in One Health: the silent pandemics	92
O.104. Report of <i>bla</i> _{NDM-5} in the Portuguese nosocomial environment: is one integron all it takes?	92
O.199. A new evolution pathway leading to azole resistance in <i>Candida glabrata</i>	94
O.213. The cell and molecular biology of carbapenem resistance in <i>Clostridioides difficile</i> .	95
O.224. From farm to fork: studying the diversity and persistence of colistin resistant <i>Klebsiella pneumoniae</i> in colistin-free chicken farms	96
O.347. Carbapenem-resistant <i>Klebsiella pneumoniae</i> can survive in surface water for at least one week	98
O.421. Analysis of vancomycin-resistant <i>Enterococcus faecium</i> causing hospital infections in Porto, Portugal, over the last decade	100
1.7 Molecular microbiology in the NGS era: early detection and identification of pathogens and its dispersion and evolution	102
O.25. Contribution of Next-Generation Sequencing to Fish Virology: virome characterization of Atlantic horse mackerels and gilthead seabreams farmed/captured along Portugal	102
O.329. Insights into the evolutionary forces shaping the pan-genome architecture of strictly clonal <i>Mycobacterium bovis</i>	104
O.444. Molecular epidemiology of Zika virus: a worldwide characterization	106
2.1 Biotechnology and Sustainability I	108
O.97. The riboflavin overproducing yeast <i>Candida membranifaciens</i> IST 626: from isolation and production optimization to genome analysis for future development of superior strains	108
O.160. Promising Genetic Determinants Of Methanol Tolerance For Yeast Robustness Engineering For Biorefineries	110
O.205. Optimizing chemical digestion approaches for polyhydroxyalkanoates extraction from biomass produced with paper-mill wastewater	112
O.233. Low-cost and eco-friendly downstream processes for single-step purification and immobilization of recombinant proteins	113
O.235. Multi-feedstock approach for ethanol production using cell surface engineered <i>Saccharomyces cerevisiae</i>	115
O.247. Impact of a DedA family protein in Indium Bioaccumulation by a <i>Rhodanobacter</i> sp. B2A1Ga4 strain	117
O.264. Engineering Ato1 transporter from <i>Saccharomyces cerevisiae</i> enables higher range of carboxylic acids uptake	118
O.405. Marine-derived actinobacteria produce bioplastics while degrading (micro)plastics?	119
2.2 Biotechnology and Sustainability II	121
O.5. <i>Zymomonas mobilis</i> : a promising microorganism for prebiotic production	121
O.38. Camelina sativa meal hydrolysate as sustainable biomass for the production of carotenoids by <i>Rhodospiridium toruloides</i>	122
O.57. Biological saline produced water treatment, a sustainable process towards lipids production	123
O.120. Microalgae granular systems feasible for the treatment of marine aquaculture streams containing florfenicol antibiotic	125

O.158. Biochemical characterization of the inner membrane cytochrome CbcL: _____	126
a gate for extracellular electron transfer in <i>Geobacter sulfurreducens</i> _____	126
O.246. Engineering a native <i>Kluyveromyces marxianus</i> transporter to reprogramme the balance between hexose and pentose transport _____	127
O.321. Unveiling the Phytochemical Nature of Acorns: The Relevance of Dehusking _____	128
O.324. Anaerobic biological removal of pharmaceuticals: impact of these micropollutants towards different microbial groups in anaerobic communities _____	129
2.3 Biotechnology and Sustainability III _____	130
O.223. Lyophilized bacterial agents for bioremediation of hydrocarbons in seawater: tackling a port oil spill accident _____	130
O.226. The Curious Case of Microorganisms and Moonmilk: _____	132
A comparative study of three non-ornated caves from the Vézère Valley (Dordogne, France) _____	132
O.270. GreenRehab: A rehabilitation system for burned soils based on the inoculation of native cyanobacteria and microalgae _____	134
O.338. Facultative anaerobic bacteria: key players in syntrophic fatty acids degradation under microaerophilic conditions _____	136
O.366. Efficient ammonium removal from marine aquaculture wastewater with microalgal-bacterial granular sludge technology _____	138
O.400. Optimization of phosphate accumulating bacterial strains for phosphorous uptake in residual waters _____	139
O.402. Dynamic columns to assess the impact of nutrients, bioaugmentation and negative pressure in bioleaching of mining residues of Panasqueira mine _____	141
O.407. Interaction of <i>Mesorhizobium quingshengii</i> J19 with high concentrations of yttrium _____	143
3.1 Microbial physiology and biochemistry: microbial activity in the context of environmental changes _____	145
O.113. <i>Erwinia</i> regulates expression of host specific virulence factors via quorum sensing _____	145
O.225. Effect of low pH in the denitrification pathway and nitrous oxide reductase _____	147
O.355. Cyanobacterial extracellular vesicles: biological functions and biotechnological applications _____	148
3.2 Biotechnology and environmental microbiology: from psychrophiles to extremophiles _____	150
O.19. Biodegradation potential of paracetamol by consortia and bacterial isolates recovered from marine organisms _____	150
O.30. Genetic characterization of viral structural-protein coding-sequences of Human polyomaviruses present in wastewater and environmental samples Lisbon _____	151
O.87. Exploring deep-sea-Actinobacteria chemical diversity by using the OSMAC approach _____	153
O.161. Light-emitting diodes effect on <i>Aspergillus</i> species in filtered surface water matrices: DNA damage, proteome response and potential reactivation _____	154
O.194. Bioprospecting Actinobacteria from poorly surveyed deep-sea habitats of the Azores and Madeira archipelagos _____	155
O.293. Impact of microbiological contamination on dark stains in granite and mural painting of religious buildings in Northern Portugal _____	157
O.386. Comparative genomics unveils fundamental traits on the biology and physiology of the bio-flavouring species of the <i>Saccharomycetaceae</i> family <i>Hanseniaspora guilliermondii</i> and <i>Saccharomyces ludwigii</i> _____	159
O.393. Occurrence and inactivation of antibiotic-resistant bacteria present in surface water _____	160
3.3 Microbiological collections: resources for a sustainable present and future _____	161
O.79. Marine host-associated bacteria show versatile secondary metabolite biosynthesis capacities and antimicrobial activities _____	161
O.323. The BACA collection as a resource for cyanobacteria diversity assessment of the Azores using a polyphasic approach _____	164
O.331. A Georeferenced Library of Native Microorganisms for Bioremediation of Oil Spills _____	166
3.4 Soil health, food security and food safety _____	168
O.174. Fresh cheese shelf-life extension at variable room temperature comparatively to refrigeration _____	168
O.178. Multilocus DNA metabarcoding of complex mixtures containing members of the Boraginaceae family for food safety and quality assurance _____	170

O.403. Surveillance study of <i>Staphylococcus aureus</i> in raw milk from Northern Portugal: enterotoxins and resistance genetic profiles _____	172
3.5 Biotech4food – nutrigenomics, personalized nutrition and food production _____	174
O.426. A sustainable alternative for cheese production – a synthetic biology approach for chymosin production using recombinant Lactic Acid Bacteria _____	174
O.438. Prevalence and control of Shiga toxin-producing <i>E. coli</i> : from diversity in dairy cattle to phage therapy ____	176
O.451. Perspectives on the development of CLA/CLNA-enriched milk through <i>in situ</i> microbial production _____	177
4.1 Teaching and Communicating in Science _____	179
O. A picture is worth a thousand words: Science Communication in Microbiology _____	179
O.94. The International Microorganism Day: an initiative from Portugal to the World _____	180
4.2 COST sessions rooms _____	181
O.417. The COST Action EuroMicroPH: Understanding and exploiting the impacts of low pH on micro-organisms_	181
O.453. YEAST4BIO COST Action: Unraveling how non-conventional yeasts can be successfully implemented in biotechnological industries for bioproducts production _____	182
5.1 Genome Editing for therapy, biotechnological & food use _____	183
O.232. New genome-editing tools for <i>Ashbya gossypii</i> _____	183
O.255. Tools provided by RNAseq for the improvement of the entomopathogenic nematode <i>H. bacteriophora</i> ____	185
O.434. Targeted removal of the <i>nth</i> nuclease gene from <i>Lactococcus lactis</i> subsp. <i>lactis</i> LMG19460 and its effects on plasmid DNA yield _____	186
5.2. Synthetic & Systems biology: experimental, theoretical and computational approaches for the analysis of micro and biological systems _____	187
O.4. A study of native tandem promoters of <i>E. coli</i> and their use in synthetic circuits _____	187
O.78. A computational approach to finding new drug targets for pathogenic <i>Candida</i> species _____	188
O.95. The N.C.Yeasttract and CommunityYeasttract databases and tools: studying transcriptional regulation in non-conventional yeasts _____	189
O.252. Study of the antibacterial mechanism of cyprosin extract against <i>Helicobacter pylori</i> cells using mid-infrared spectroscopy and chemometrics methods _____	191
O.396. Prospecting new-to-nature pathways to implement microbial production of levulinic acid _____	193
5.3 New approaches for disease control and treatment _____	195
O.209. Mycobacteriophage Ms6 LysB: a lipolytic lysis enzyme with Peptidoglycan Binding Affinity _____	195
O.210. IR Bactyping, the next cornerstone for <i>Klebsiella pneumoniae</i> strain typing towards real-time infection control _____	197
O.248. A genome-wide level perspective on the microevolutionary trajectory by endemic <i>Mycobacterium tuberculosis</i> strains in Portugal: impact for phenotypic resistance levels and therapeutic efficacy _____	199
O.275. Rifabutin liposomes: an effective drug delivery therapy for biofilm associated <i>S. aureus</i> infections _____	201
O.394. Targeting diabetic foot infection through a novel biomaterial-based topical therapeutic approach _____	203
O.432. <i>Helicobacter pylori</i> prophages: screening, detection, induction and potential therapeutic use _____	205
5.4 Digital Data, Big Data & AI in Microbiotech _____	207
O.85. Learning from fluorescence: online multiparameter monitoring of a microalgae culture _____	207
O.188. eHooke: a framework for automated image analysis of <i>Staphylococcus aureus</i> based on cell cycle progression _____	209
Poster Presentations _____	211
1.1 Zoonoses, emerging and re-emerging diseases: virology, bacteriology, mycology and parasitology _	211
28. Biosafety measures in freshwater fish ex situ breeding programs - a preliminary approach _____	211
29. Influence of climate change on growth, biofilm production and antimicrobial resistance profiles in <i>Aeromonas</i> spp. _____	213
45. Medium matters: influence of different mediums and atmospheres on the haemolytic profiles of canine oral enterococci _____	214
53. Evaluation of <i>Aliarcobacter butzleri</i> response to bile salts _____	215

55. Effect of the application of organic acids on the spoilage of bovine viscera for human consumption	216
56. Evaluation of citric and lactic acid application in the behavior of <i>Listeria monocytogenes</i> on pig carcasses samples	218
81. Presence of viable pathogenic leptospire in environmental samples from three districts of mainland Portugal: a pilot study.	220
109. Canine and feline haemoparasites – brief report of the veterinary hospital.	221
153. Tracking Tick-borne rickettsiae in a national park in Portugal	222
185. Characterization of Hazara virus infection in <i>Hyalomma lusitanicum</i> tick cell line.	224
193. Microbicidal activity of macrophages exposed to pathogenic spirochetes <i>Borrelia garinii</i> and <i>B. lusitaniae</i> species	225
195. Deciphering the virulence potential of human isolates of <i>Aliarcobacter butzleri</i>	227
198. Optimization of a Real-Time PCR protocol using primers based on the <i>lipL32</i> gene for <i>Leptospira</i> DNA detection in biological and environmental samples	228
222. The <i>in-vitro</i> antidermatophytic activity of <i>Withania Chevalieri</i> an endemic plant from Cape Verde	229
281. The role of wild animals and associated ticks in <i>Coxiella burnetii</i> natural cycle	231
376. Proposing a new <i>Shewanella</i> species causing human infections	233
401. Development of a standard operation procedure (SOP) for the polyphasic identification of clinically relevant species	234
452. Development of new diagnostic tools for the detection of <i>Toxoplasma gondii</i> .	236
1.2 Microbiome in Health & Disease: microbial diversity and host-microbes interactions	237
10. Evaluation of methicillin-resistant <i>Staphylococcus aureus</i> carriage in the elderly in Portugal using selective enrichment followed by qPCR	237
11. Induction of multinucleation in infected host cells by a <i>Chlamydia trachomatis</i> inclusion membrane protein	238
12. Intra-species interactions in pneumococcal biofilms	239
65. Functional and comparative metagenomics of viral communities in healthy and diseased octocoral tissue and seawater	240
86. The <i>Roseibium album</i> (<i>Labrenzia alba</i>) pangenome reveals multiple symbiosis factors likely underpinning host-microbe relationships in the marine benthos	243
88. Phylogenetic and functional analysis of orthologs of CteG, a <i>Chlamydia trachomatis</i> effector protein	246
89. Bacterial Vaginosis multi-species biofilms: can standard quantification methods accurately quantify <i>in vitro</i> biofilms?	247
93. Identification of <i>Clostridioides difficile</i> biofilm matrix proteins	248
99. Incl is a novel <i>Chlamydia trachomatis</i> protein interacting with host cell lipid droplets and eukaryotic 14-3-3 proteins	249
101. Study of the interaction between vaginal and intestinal lactobacilli, <i>Candida albicans</i> and <i>Candida glabrata</i>	250
103. Insights into the adaptation of <i>S. epidermidis</i> biofilms to effectors of the immune system	251
122. Characterization of single- and multi-species Bacterial Vaginosis (BV)-associated biofilms in an ex vivo 3D human vaginal epithelium model	252
124. The role of the gene <i>codY</i> in <i>S. epidermidis</i> biofilm formation	253
140. The putative toxin-antitoxin <i>mazEF</i> gene cluster is not involved in cell death in <i>S. Epidermidis</i>	254
148. Identification of amino acid residues important for the subcellular localization of the <i>Chlamydia trachomatis</i> effector CteG in host cells	255
150. Lytic exit of <i>Chlamydia trachomatis</i> from host cells is mediated by the CteG effector protein	256
180. Evaluation of mannans from <i>Saccharomyces cerevisiae</i> as potential human microbiota modulators	257
190. Molecular players involved in the transition of free-cells to multicellular aggregates in <i>Burkholderia multivorans</i>	258
191. Peptide-rich extracts from spent yeast as potential microbiota modulators	260
197. Oral poliomyelitis revaccination is associated with changes in gut and upper respiratory microbiomes of infants	261
204. Nanopore sequencing approaches to profile the gut microbiota of mice infants exposed to alcohol in utero	262
208. <i>Staphylococcus epidermidis</i> molecular responses to pH changes during human infection – a multiomics approach	264
229. Comparative metabolomic study indicates a metabolic re-route of a pathogenic <i>Staphylococcus epidermidis</i> strain at blood pH	266

231. Impact of a pulse-based vegetarian lunch meal on the modulation of gut bacterial sub-populations	268
268. Role of ScwdABC system of <i>Staphylococcus aureus</i> in biofilm formation and infection	270
286. The microbiome(s) of Senegalese sole fed with different diets	272
294. Mass Spectrometry – Based proteomic and metabolomic profiling of serum samples for discovery and validation of TB diagnostic biomarker	274
295. Promoting a healthy microbiome with <i>Lactobacillus</i> isolated from human milk	276
300. Uncovering the commensalism vs pathogenicity in <i>Staphylococcus Epidermidis</i> – A combined genomic/proteomic approach	277
303. Two novel <i>Gardnerella</i> species isolated from urine of women	279
320. Chronic vulvovaginal <i>Candida</i> spp clinical isolates are more resistant to phagocytosis in in-vitro models of infection	281
372. Do all prebiotics have the same impact on my gut microbiota? A case study using an <i>in vitro</i> gastrointestinal model to assess functional oligosaccharides added to milk	282
375. Identification of new players of lysozyme resistance in <i>Staphylococcus aureus</i>	283
383. Peptidoglycan amidation impact on biofilm formation - role of eDNA and Atl-	285
389. Establishment of the MurT-GatD complex as a new virulence factor	286
427. The genetic basis of bacterial adaptation to a complex environment and its consequences on <i>Caenorhabditis elegans</i> ability to respond to environmental and life-history changes.	287
446. Characterization of small noncoding RNAs expressed by <i>Burkholderia cenocepacia</i> when infecting <i>Caenorhabditis elegans</i>	288
465. Probiotic adhesion to skin keratinocytes and underlying mechanisms	290
1.3 Microbiology & Biotechnology in COVID-19: new approaches for new diseases	291
14. Preventing COVID-19 outbreaks through active screening of civil servants - A partnership between local government, a primary-care center and a research institute	291
145. SARS-CoV-2 contamination potential in environmental and wastewaters in the Algarve region, Southern Portugal	293
165. Repositioning drugs with potential activity against SARS-CoV-2 infection	295
263. Evaluation of minimum inhibitory concentrations of plants extracts against several bacterial pathogens	297
278. Experimental approach to detect SARS-CoV-2 in aerosol suspensions in COVID19 patients domiciliary setting	298
311. UVC radiation and Ozone as disinfection methods against a viral surrogate of SARS-CoV-2	300
315. Antioxidant activity, phenolic profile, cytotoxicity and genotoxicity of plant extracts	302
322. Ozone and UVC radiation as disinfection strategies of textile substrates	303
341. Predominant phenolic compounds in Mountain plants with possible antiviral potential against SARS-CoV2	304
449. Assessing the microbiological efficacy of three scalable decontamination methods as a strategy to address the need for respiratory protective devices during a pandemic	306
1.4 & 1.6 Biopharmaceuticals & Vaccines: design, analytical methods and bioprocessing opportunities	308
13. An alternative way to purify recombinant E6 protein from <i>Escherichia coli</i> cells: a target for cervical cancer	308
27. QAS-hydrogels as a promising microbicide against sexually and perinatal transmitted infections	309
152. Surfomics reveal a new target for the development of immunotherapies to tackle <i>Burkholderia cenocepacia</i> infections	310
302. A chromatographic strategy for the purification of small heat shock protein nanocages	311
364. Recombinant CBM-fusion technology towards production of novel cellulose-based materials with biomedical interest	312
379. Bacterial cellulose for burn wounds dressings	314
385. Essential Oils of <i>Mentha pulegium</i> , <i>Mentha spicata</i> and <i>Calamintha nepeta</i> , from Alentejo: Antimicrobial Activity against Biodeteriogenic Microorganisms and Immobilization in β -cyclodextrin Complexes	315
392. Development of antimicrobial polymeric surfaces	317
413. Lentiviral vector components expression levels and stability: impact on constitutive producer cell yields and vector quality	319
422. Establishing new cell platforms for scalable lentiviral vector production	320
424. Characterization of a novel ferulic acid pathway in <i>Lactococcus lactis</i>	321
1.5 Antimicrobial Resistance in One Health: the silent pandemics	323
16. Heteromeric phage endolysins - not just an eccentricity of few	323

18. Determinants of bacterial tolerance to endolysin-based enzybiotics _____	325
26. Antibiotic Resistance Mechanisms in Environmental and Pathogenic Carbapenem Resistant Bacteria from a Portuguese Wastewater Treatment Plant _____	326
35. A novel target in TB-therapy: the <i>N</i> -glycolylation of mycobacterial peptidoglycan contributes to antibiotic resistance and intracellular survival _____	328
36. Finding Key Genomic Markers of Beta-lactam Susceptibility in Clinical Isolates of <i>Mycobacterium tuberculosis</i> _____	329
43. Ascertaining mycobacterial inducible resistance _____	331
to peptidoglycan synthesis inhibitors _____	331
46. Antimicrobial resistance and virulence profiles of Enterobacteriaceae isolated from two-finger and three-finger sloths (<i>Choloepus hoffmanni</i> and <i>Bradypus variegatus</i>) of Costa Rica _____	332
52. Evaluation of efflux activity in <i>Staphylococcus pseudintermedius</i> and its contribution for biocide decreased susceptibility _____	333
54. Biofilm production by <i>Staphylococcus pseudintermedius</i> associated with animal pyoderma _____	335
70. Unravelling the drivers of antimicrobial resistance in <i>Staphylococcus aureus</i> from wildlife _____	337
71. In vitro interactions during bacterial vaginosis development demonstrate that multi-species biofilms have enhanced antimicrobial tolerance _____	338
82. Insights into pESI-like megaplasmid of <i>Salmonella</i> Infantis ST32 spreading in the poultry industry _____	339
83. Overview on antimicrobial resistance of <i>Salmonella</i> enteritidis ST11 in the poultry industry, Portugal 2014-2020 _____	340
84. Overview on the resistance to critically important antibiotics (CIA) from food-producing animals and food _____	341
90. Antibacterial and antiviral activity of Preyssler polyoxotungstate P ₅ W ₃₀ _____	342
107. Bacterial urinary tract infections in pets: diversity vs antibiotic resistance and its significance for public health. _____	343
126. Freshwater cyanobacteria: a new player in the One Health puzzle _____	344
206. Detection of β -lactamases and the simultaneous presence of increased efflux activity in <i>Enterobacteriaceae</i> using a 96-well microplate method _____	346
212. The genetic circuitry controlling production of a MecA-like penicillin binding protein required for carbapenem resistance in <i>Clostridioides difficile</i> _____	348
214. Bacterial Peroxidases from Pathogenic Bacteria _____	349
221. Twort CHAP domain secondary binding site specifically targets the peptide component of <i>Staphylococcus aureus</i> peptidoglycan _____	350
228. The dual binding mode of Twort SH3b domain to <i>Staphylococcus aureus</i> peptidoglycan _____	351
236. Evaluation of the genetics underlying a putative acquisition of antimicrobial resistance to novel alkylaminophenols _____	353
237. Are quaternary ammonium surfactants a good prophylactic option against <i>Streptococcus agalactiae</i> vertical transmission? _____	355
242. The pig production chain in Portugal is a key reservoir of antibiotic-resistant and pathogenic Enterobacteriaceae _____	356
243. Assessing the role of phenotypic heterogeneity on <i>Staphylococcus aureus</i> tolerance to β -lactams _____	358
244. Screening of stress effect on the survival of clinical and environmental antibiotic resistant <i>Escherichia coli</i> and <i>Klebsiella pneumoniae</i> isolates _____	359
245. Conjugation of imidazolium-based ionic liquids to antimicrobial peptides: effects on antimicrobial activity and cytotoxicity _____	361
249. Virulence traits of <i>Staphylococcus aureus</i> associated with skin and soft tissue infections in humans and animals _____	362
250. Co-carriage of clinically important antimicrobial resistance genes by healthy companion animals and their human household members _____	364
259. Detoxification of toxic intermediates produced by oxygen reduction in human pathogens _____	366
266. Evaluation of IR Biotyper for long-term surveillance of carbapenemase-producing <i>Klebsiella pneumoniae</i> clinical isolates _____	368
298. A hygiene and monitorization program for small animal veterinary practices – A Pilot-study _____	370
299. ESBL and AmpC producing Enterobacteriales in urban waters of Douro River in Portugal – _____	372
Is the environment a mirror of community intestinal colonization? _____	372

301. Characterization of bactericidal activity of new NHC carbenes and their Cu(I) and Ag(I) complexes against pathogenic bacteria	374
325. Optimized Red Nile semi-automated fluorometric method shows competition between substrates of the AcrAB-TolC efflux pump of <i>Escherichia coli</i>	376
326. Measuring efflux activity in <i>Escherichia coli</i> using Hoechst 33258, a bis-benzimidazole derivative	378
332. The role of the pork production chain as a reservoir and transmission route of Enterobacteriaceae resistant to last resort antibiotics	380
344. Using genomics to uncover the role of SMR efflux pumps to <i>Acinetobacter baumannii</i> drug resistance	382
349. Comparative genomics insights into clinical or environmental methicillin-resistant <i>Staphylococcus aureus</i>	384
359. Exploring <i>Enterococcus</i> susceptibility to co-occurring stresses in the poultry meat production chain: antibiotics, copper, acidic pH and peracetic acid	386
365. Molecular characterization of <i>Staphylococcus pseudintermedius</i> associated with skin and soft tissue infections in pets, Portugal	388
367. Exotic Pets as a reservoir of MultiDrug Resistant priority pathogens with special focus on <i>Enterococcus faecalis</i> (2017-2019 Toulouse, France)	390
369. Pan-genome and pan-resistome analysis of bla _{VIM-2+} or bla _{NDM-1+} Carbapenem resistant <i>Pseudomonas aeruginosa</i>	392
370. Quaternary Ammonium Compounds' activity and occurrence of tolerance associated genes among <i>Enterococcus faecalis</i> and <i>Enterococcus faecium</i> from diverse origins and timespans	394
374. Antibacterial and synergetic activity of essential oils of <i>Thymus capitellatus</i> , <i>Origanum virens</i> and <i>Origanum macrostachyum</i> of Alentejo (Portugal)	396
377. Assessment of MICs for <i>Mycobacterium tuberculosis</i> complex clinical strains using the EUCAST reference method	398
381. Emergence of staphylococcal non-wild-type populations to biocides in a One Health perspective: are there reasons for concern?	400
387. New CRISPRi System to Inhibit Transcription of Essential Genes in <i>Staphylococcus aureus</i>	402
397. Exploring environmental sources and transmission routes of multidrug-resistant <i>Escherichia coli</i> in Lettuce Greenhouses	403
404. Antimicrobial resistance and molecular typing of <i>Staphylococcus aureus</i> causing bacteraemia in children aged less than 5 years in Southern Mozambique, 2001-2019	405
416. Fluoroquinolone resistance and associated mechanisms in <i>Staphylococcus pseudintermedius</i>	407
420. <i>Candida glabrata</i> response to the antifungal fluconazole is partially coordinated by the transcription factor Zap1	409
435. Microemulsions based on deep eutectic solvents to tackle resistant ocular infections	410
450. Raw pet food: new route for bacterial zoonotic health risks spread in Europe	412
455. Irrigation Water and Soil as sources of Pathogenic and Antibiotic Resistant Bacteria in Portuguese tomato greenhouses	414
459. Antimicrobial screening of <i>Lavandula</i> spp. hydrodistillation products against potentially pathogenic bacteria	416
1.7 Molecular microbiology in the NGS era: early detection and identification of pathogens and its dispersion and evolution	418
9. Pneumococcal carriage among Portuguese children after the introduction of the 13-valent pneumococcal conjugate vaccine (PCV13) in the National Immunization Plan	418
112. Vancomycin resistant <i>Enterococcus faecium</i> causing infection in portuguese hospitals: predominance of the ST17- <i>vanA</i> lineage	420
121. Pretreatment of biological samples using three-phase partitioning systems: a tool to enhance early diagnosis	422
170. Genome analysis of <i>Diaporthe amygdali</i> and <i>D. eres</i> : unraveling adaptation for plant cell wall degradation and its pathogenicity potential	423
200. Unveiling the genomic basis for differential biofilm formation capacity in <i>Candida glabrata</i> clinical isolates	425
423. Genes contributing to high-level oxacillin resistance among MRSA isolates carrying different methicillin genetic determinants	426
2.1 Biotechnology and Sustainability I	427

7. Assessing predatory activity of native Portuguese filamentous fungi against coccidia from peacocks (<i>Pavo cristatus</i>) – <i>in vitro</i> preliminary results	427
40. Improving the production of polymers using biocatalysis	428
41. Design of an integrated platform based on aqueous biphasic systems for the enzymatic production of polydopamine and biocatalyst reuse	430
58. Obtention of PHB from cocoa shell wastes by fermentation	432
74. Bacterial Cellulose: From Biotechnology to Bio-Economy	433
76. Sugarcane-derived biogenic silica particles as a topical delivery system for retinoids	434
96. Enhancing anaerobic digestion of lignocellulose biomass using a thermophilic microbial consortium pretreatment	435
105. Aerobic granular sludge, a feasible technology for the treatment and recirculation of trout aquaculture water streams	436
115. Production of 3D biocomposite scaffolds based on polyhydroxyalkanoates (PHAs) and FucoPol	437
117. Coupling microbial engineering with structural biology: a new foundation to solve <i>Geobacter's</i> extracellular electron transfer paradigm	439
130. Assessment of bioethanol production from <i>Eucalyptus globulus</i> bark kraft pulp	441
143. Characterization of the thermostable bacterial biosurfactant produced by <i>Burkholderia thailandensis</i>	442
155. Sugarcane as a sustainable source of biologically active carboxymethyl cellulose	443
156. Sugarcane bagasse cello-oligosaccharides as prebiotic agents	444
171. Integrated genomic and metabolomic analysis of the marine fungus <i>Emericellopsis cladophorae</i> : insights into saltwater adaptability mechanisms and its biosynthetic potential	445
175. Functional screening for the detection of β -glucosidase activity in a metagenomic library obtained from a compost sample	446
176. Engineering redox proteins for optimal bioenergy production	448
184. Redirecting the reducing power in <i>Starmerella bombicola</i> towards production of compounds with industrial interest	449
196. Development of GABA-enriched fermented beverage based on sweet whey	450
211. Microwave-assisted extraction strategies for biologically active molecules from <i>Tetraselmis chuii</i> and <i>Chlorella vulgaris</i>	451
216. Phenolic compound modulation in yeast fed-batch fermentation using sugarcane syrup as feedstock	452
238. Microalgal biomass production in a raceway system using meat processing water as feedstock	454
239. Yeast platform for the production of 5-hydroxymethylfurfural-derivatives: <i>Saccharomyces cerevisiae</i> as a whole cell biocatalyst	455
241. Project BIOREM: Bioremediation of hydrocarbon pollutants by autochthonous microorganisms in aquatic environment	456
251. Overexpression of bacterial metallothioneins increases the Gallium and Indium recovery	458
254. Metal biosorption by bacterial biopolymers	460
261. Comparative genomics of bacterial mine isolates to unravel metabolic features linked to gallium bioleaching ability	462
267. The importance of QrcABCD-QmoABC redox-loop for H_2 /formate-sulfate respiration in sulfate reducers	464
269. Effect of extraction methods on antioxidant activities of <i>Tetraselmis chuii</i>	465
274. Improving <i>Desulfovibrio vulgaris</i> Hildenborough [NiFeSe] Hydrogenase Resistance to Oxidative Inactivation	466
277. Recombinant expression and insecticidal activity of one ShK domain of the entomopathogenic nematode <i>Steinernema carpocapsae</i> in fusion with –DsbC	467
282. Application of a fungal extract with laccase activity to improve the enzymatic hydrolysis of eucalyptus bark residues	468
283. Development of an electrotransformation protocol for the novel acetogen <i>Acebotacterium wieringae</i> strain JM	470
284. Targeting Aerobic Granular Sludge Microbiome Salt Adaptation	472
285. Exploring the potential of primary sludges from paper mills for the production of value-added compounds	473
291. Kudzu starch structure, physical, and mechanical properties, a starch with potential application on cultural heritage conservation	475
292. Soil microbiota benefits from phytoremediation coupled to metal-resistant rhizobacteria	477

297. Application of bacteriocin-producer in the reduction of cheese maturation time _____	478
304. Over-expression of AMPs in industrial <i>Saccharomyces cerevisiae</i> strains as a biocontrol approach for fuel ethanol fermentations: cloning strategy and characterization _____	479
305. Antibacterial biocomposites of a genetically engineered protein polymer incorporating essential oil from <i>Mentha piperita</i> _____	481
310. Wine industry residues for biotechnological production of xylitol _____	483
313. Impact of gastrointestinal digestion on the biological activities of new functional cereal-based granules based on by-products from the food industry _____	484
314. Anaerobic granular sludge as biocatalyst for CH ₄ production at moderate H ₂ /CO ₂ pressures _____	486
316. Unraveling the Role of Dehulling on Acorn's Nutritional Composition _____	488
319. Production of recombinant bioactive antimicrobial peptides using a thermoresponsive fusion tag _____	489
334. Development of efficient industrial xylose-fermenting <i>Saccharomyces cerevisiae</i> strains using genes from native xylose-utilizing yeasts _____	491
335. Xylulokinases from two native xylose-fermenting yeasts improve xylose conversion in recombinant <i>Saccharomyces cerevisiae</i> _____	492
351. Antimicrobial potential of formulations, incorporating spent yeast derived from synthetic biotechnology, against <i>Pseudomonas spp.</i> _____	493
353. Valorization of biotechnology derived spent yeast as potential ruminant feed additive _____	495
356. Growth of <i>Saccharomyces Cerevisiae</i> : Screening the Best Conditions _____	496
357. Biosolvents, ionic liquids, or eutectic solvents? In the search of best solvents for the extraction of carotenoids from yeast cells _____	497
362. Resveratrol production by recombinant robust yeast strains for valorisation of wine waste _____	499
368. Lignin-based adhesives as a sustainable strategy for conservation of cultural goods _____	501
371. High value tellurium nanoparticle formation by <i>Paenibacillus pabuli</i> _____	503
395. Dye decolorization by yeasts: towards an innovative solution for textile effluents _____	505
409. Optimization of hemicellulose extraction from sugarcane straw by autohydrolysis _____	507
410. Screening of antimicrobial activity in fractionated synthetic biology industry waste streams _____	508
411. Synthetic biology byproducts valorization into fish-feed prebiotic ingredients _____	510
425. Bacterial Biodegradable Polymers in a Circular and Sustainable Bioeconomy _____	511
428. Microalgae-bacterial granular sludge for the treatment of low carbon and nutrient loaded effluents _____	513
430. Pre-treatment of sugarcane straw and bagasse with ionic liquid for the production of cellulosic ethanol and lignin _____	515
431. Bacterial cellulose production from terephthalic acid and ethylene glycol by <i>Komagataeibacter xylinus</i> DSM 2004 and 46604 _____	517
437. Genome Sequence and Annotation of <i>Lactococcus lactis</i> Strain LMG 19460 and Novel Strains IST-LL01 and IST-LL02 _____	518
440. Fertilizer technology- a new path for the valorisation of horticultural residues _____	520
442. Process intensification for the recovery of intracellular carotenoids from <i>Rhodotorula glutinis</i> yeast _____	521
448. Potential of sugarcane straw extracts as preservatives for the cosmetic industry _____	523
454. Overgrowth control of potentially hazardous bacteria in ozonated wastewater through competition: microbiome analysis _____	524
457. Sugarcane bagasse as source of bioactive lignin: influence of pretreatment on the antioxidant and antibacterial activities _____	526
460. Coatings of chitosan and chitosan with essential oil of <i>L. stoechas</i> subsp. <i>luisieri</i> for the control of spoilage microorganisms in post-harvest conservation of <i>Prunus avium</i> fruits _____	528
464. The effect of calcium concentration on culture medium for the production of biosandstones _____	530
2.2 Biotechnology and Sustainability II _____	531
102. Sustainable lysis of <i>Bacillus subtilis</i> biomass to recover the biopharmaceutical L-asparaginase _____	531
230. Ionic liquids as promising solvents for biocatalysis _____	533
2.3 Biotechnology and Sustainability III _____	534
192. Mining seaweed-associated Actinobacteria for biotechnologically-relevant compound discovery _____	534
215. Membrane transporters in the bioproduction of organic acids: state of the art and future perspectives for industrial applications _____	536

218. Expanding the knowledge on the skillful yeast <i>Cyberlindnera jadinii</i>	537
240. Diversity and hydrocarbon-degrading potential of deep-sea microbial community from the Mid-Atlantic Ridge, south of the Azores (North Atlantic Ocean)	538
258. <i>Streptomyces</i> spp. rRNA-FISH probe to apply in cultural heritage materials: <i>in silico</i> design and experimental evaluation of specificity and performance	540
262. Uncovering novel <i>Cyberlindnera jadinii</i> carboxylate transporters for the improvement of microbial cell factories	542
279. Unraveling the diversity and biotechnological potential of ascidian-associated Actinobacteria	543
287. Looking into marine macroalgae associated bacteria to detect the presence of CAZymes	545
330. Biotechnological exploitation of <i>Torulaspota delbrueckii</i> through phenotypic and metabolic characterization	546
348. Diversity of metallophytes and metal(loid)-tolerant bacterial strains in a portuguese mine for phytotechnologies purposes	548
360. Biodiversity assessment in Marine caves of the Algarve coast and: anti-tumoral prospection of bioactive compounds produced by bacterial isolates	550
418. Potential of native microorganisms to degrade two halogenated pharmaceuticals	551
419. Ectomycorrhizal fungi inocula optimization to aid the health status of trees in the everchanging environment of cities	553
436. Comparative analysis of microbial communities of two craft beer styles	554
447. Antifungal properties of crude extracts from deep-sea- and macroalgae-associated Actinobacteria	557
3.1 Microbial physiology and biochemistry: microbial activity in the context of environmental changes	558
34. Potential of marine bacteria to biodegrade conventional and bio-based microplastics	558
60. Detection and cellular viability assessment of <i>Erwinia amylovora</i> in environmental samples from pear cv. "Rocha" orchards by immuno-flow cytometry	559
91. Haem biosynthesis in Gram-negative pathogens	561
159. Response to acetic acid stress in yeast: focus on cell wall remodeling	562
183. Production and purification of substrates for the characterization of phosphatidylinositol mannoside biosynthesis in <i>Mycobacterium tuberculosis</i>	564
186. The haem biosynthesis pathway of <i>Campylobacter jejuni</i>	565
276. Investigating how exoribonucleases and small RNAs influence	566
318. Extracellular vesicles biogenesis in cyanobacteria: towards the identification of underlying mechanisms	567
398. Impact of Climate Change on Ochratoxigenic Fungi	568
458. Enzymes and secondary metabolites profiles of <i>Gnomoniopsis smithogilvyi</i> are affected by chestnut medium	569
3.2 Biotechnology and environmental microbiology: from psychrophiles to extremophiles	571
20. Enrichment, isolation and characterization of acidophilic sulfate reducing bacteria from Sao Domingos mine	571
39. Bacterial food spoilage events: the first complete genome sequence of the off-flavour producer <i>Alicyclobacillus acidoterrestris</i> DSM 3922 ^T	573
51. Fungal treatment of Kraft Black Liquor using <i>Phanerochaete chrysosporium</i> and <i>Aspergillus uvarum</i>	575
63. Profiling the volatile exometabolome of <i>Pedobacter lusitanus</i> NL19	576
75. BTEX removal by <i>Aspergillus niger</i> in saline medium	577
227. Biodiversity assessment in Palaeolithic caves: the case of Escoural cave	578
260. Molecular characterization and antimicrobial properties of bioactive compounds produced by bacterial isolates from underwater caves of the Algarve coast	579
306. Recovery of heterotrophic bacteria from alkaline mine residues and their capacity to leach impurities from secondary sources of raw materials	581
312. Preliminary study on the usage of marination against common food-borne pathogens present in poultry meat.	583
340. Spatial-temporal dynamics of <i>Vibrio</i> spp. across an urban temperate estuary (River Douro estuary, NW Portugal)	585
345. Microbiome-metabolome relations in the halophyte <i>Salicornia ramosissima</i>	587
3.3 Microbiological collections: resources for a sustainable present and future	589
23. Bioprospecting bacterial symbionts of marine animals: sponge and coral-associated <i>Gammaproteobacteria</i> inhibit growth of notorious human and aquaculture pathogens	589

31. Antimicrobial peptides of <i>Saccharomyces cerevisiae</i> – a proteomic study	592
119. Microalgae: sustainable sources of nutrients	594
127. Estela Sousa e Silva Algae Culture Collection (ESSACC) – a resource of cyanobacterial strains	595
220. Characterization of the antagonistic potential of wine yeasts for biocontrol of phytopathogenic fungi in grapes	597
380. Isolation and phylogenetic identification of marine sponge-associated Actinobacteria from Mozambique and Portugal	599
382. Biotropical Resources (GHTM/IHMT NOVA biobank): Biobanking African Pediatric Upper Respiratory and Gut Microbiomes	601
441. The importance of a biological resources center in the sustainable development of a low-density region	603
445. <i>Faunomonas pinastri</i> gen. nov., sp. nov., an endophyte from a pine tree of the class <i>Alphaproteobacteria</i>	605
3.4 Soil health, food security and food safety	607
32. Lactic acid bacteria from Azeitão and Nisa PDO-cheeses: evaluation of technological and probiotic potential	607
33. <i>Enterococcus</i> spp. from Azeitão and Nisa PDO-cheeses: surveillance for antimicrobial drug resistance	608
50. Inactivation of <i>Listeria Monocytogenes</i> in Milk by Pulsed Electric Field (PEF) and Mild Heating	609
100. Phage Therapy for the Inactivation of <i>Aeromonas hydrophila</i> in Cockles During Depuration: <i>In Vitro</i> and <i>In Vivo</i> Preliminary Studies.	611
108. Shelf-life extension of squid and shrimp skewers through the application of vinegar solutions	613
154. Cellulose from sugarcane bagasse as a potential prebiotic agent	615
164. Light emitting diodes as an alternative disinfection method to inactivate bacteria associated with ready-to-eat salads	617
166. Microbial and sensory evaluation of hyperbaric storage at room temperature of a commercial ready-to-eat fish soup	618
168. Combination of moderated high pressure and ultrasound treatments for subsequent liquid whole egg improved thermal pasteurization	620
173. <i>Byssochlamys nivea</i> ascospore germination control by hyperbaric storage – dependence of the activation procedure	622
177. Molecular detection of species from the Boraginaceae family to safeguard food and feed safety, integrity, authenticity, and quality	624
179. Effect of <i>Thymus zygis</i> essential oil against <i>Listeria monocytogenes</i> and its application on food	626
265. Determination of the minimum inhibitory concentrations of the antibacterial oregano and thyme essential oils in the presence of polysorbate 80	628
307. Rainfall impact on soils subjected to no-tillage and crop-livestock integration treatments: insights on the microbial community succession following drought or flood using PLFA	629
327. Optimization of an innovative phagomagnetic protocol towards a selective and rapid <i>Listeria monocytogenes</i> detection	631
333. Exploring the mechanism of action of cymoxanil	633
336. Elderberries: Extraction and Application in Yogurt	634
337. Preparation and characterization of bioactive chitosan-based films incorporated with olive leaves extract for food packaging applications	635
352. Microbial growth on cheese surface covered with propolis	637
373. Integrating cutting-edge technologies to evaluate the effects of high-pressure processing on sea bass fillets quality and safety	638
390. Comparison of different preservation methods on the microbiological, texture and color properties of industrial bread during storage time	640
391. <i>Penicillium crustosum</i> as a potential OTA producer – new insights from whole-genome sequencing of strain MUM 16.125	642
399. Nitrite in meat products: alternatives from natural sources	644
443. Bioactive alternatives to mitigate foodborne pathogens	646
456. Are nitrate/nitrite free dry-cured sausages microbiologically safe?	647
475. Microalgae biomass as filler for starch films toward active and sustainable food packaging	648
476. Developing DNA-based methods to detect the presence of adulterant milk in Serra da Estrela Cheese	650
3.5 Biotech4food – Nutrigenomics, Personalized Nutrition and Food Production	652

106. Comparison of dietary fiber content of carob (<i>Ceratonia siliqua</i> L.) flour as related to seed presence and roasting effects _____	652
271. Effect of High Pressure Processing on Microbial Quality, Physicochemical Quality and Antioxidant Properties of Rosemary Honey During 2 Years of Storage _____	654
328. Protective effect of <i>Chlorella vulgaris</i> on potential probiotic lactic acid bacteria during in vitro gastrointestinal digestion _____	656
339. Influence of modified atmosphere packaging on multicereal bread quality and microbiological stability over storage _____	657
350. Optimization of Elderberry Fermentation: Screening Six _____	658
358. Impact of vegetable oils on Lactocaseibacillus rhamnosus and Bifidobacterium animalis subsp. lactis growth _____	660
363. <i>Fucus vesiculosus</i> and <i>Porphyra dioica</i> Seaweeds Cultivated in Fully Controlled Closed Systems: Proximate and Elemental Composition and Antioxidant Potential _____	661
429. Metagenomic characterization of bacterial and yeast communities in donkey milk fermented with kefir grains _____	663
461. Chemical Characterization of <i>Tetraselmis chuii</i> : _____	664
4.1 Teaching and Communicating in Science _____	666
342. A Microbiology Outreach Project Within Training in Responsible Research And Innovation _____	666
378. MicroMundo/Tiny Earth: an international citizen science project to fight antimicrobial resistance from universities to Basic/high school's academic communities from Center region of Portugal. _____	668
5.1 Genome Editing for therapy, biotechnological & food use _____	670
151. Isolation of novel <i>Chlorella vulgaris</i> mutants with higher biomass productivities for food and feed applications: a non-GMO strain improvement approach _____	670
384. Screening of Lactic Acid Bacteria strains for Electrotransformation Efficiency _____	672
5.2 Synthetic & Systems biology: experimental, theoretical and computational approaches for the analysis of micro and biological systems _____	673
111. Predictive oxidative evaluation of lipid samples using Differential Scanning Calorimetry _____	673
414. Reconstruction of the Metabolic Network of Lactococcus lactis strains IL1403 and LMG 19460 _____	674
5.3 New approaches for disease control and treatment _____	676
8. An alternative approach to blood disinfection: photoinactivation of bacteria, fungi and viruses _____	676
44. Antimicrobial activity of MgO nanoparticles against <i>Enterococcus faecalis</i> using an <i>ex vivo</i> dental model _____	677
47. Pexiganan-nisin biogel in combination with conventional antibiotics against polymicrobial diabetic foot infections _____	678
64. New infection models to test drugs against nontuberculous mycobacteria _____	679
98. Bacteriophages for biocontrol of <i>Pseudomonas syringae</i> pv. <i>actinidiae</i> : Use of single phage suspensions and phage cocktails _____	681
116. Abrogation of endocytosis blocks the spread of alpha-synuclein in the yeast model of synucleinopathies _____	682
118. The effect of alternative sugars in the yeast <i>Saccharomyces cerevisiae</i> fitness _____	683
123. Resazurin Cell Viability Assay: an Effective Protocol to Assess Bacterial Inactivation by Phages _____	684
125. Phages for food pathogens inactivation: preliminary results in milk _____	685
133. Identification of new drugs for <i>Neisseria gonorrhoeae</i> using an <i>in silico</i> repurposing strategy _____	686
135. Influence of the water physicochemical parameters in the antimicrobial photodynamic therapy effectiveness _____	688
169. Pyranoanthocyanins affecting the expression of Quorum Sensing-related genes in <i>Pseudomonas aeruginosa</i> and <i>Staphylococcus aureus</i> biofilms _____	689
187. Carbon Monoxide Releasing Molecules and infectious diseases _____	690
219. Disinfection of textiles and footwear using an ultrasonic nebulizer cabinet _____	692
253. Repurposing of new drug candidates for Buruli ulcer treatment _____	694
288. Preliminary characterization of a novel antimicrobial surface coating against significant pathogens _____	695
290. Developing a handheld biosensor to detect <i>Staphylococcus aureus</i> infections _____	696
308. Marine Drug-Like Leads for SARS-CoV-2 Main Protease Inhibition Predicted by a Computer-Aided Drug Design Approach _____	698
317. Biocontrol of <i>Pseudomonas syringae</i> pv. <i>actinidiae</i> in kiwi plants using bacteriophages _____	700

346. Optimizing spray-drying conditions for encapsulation of the next generation probiotic <i>Akkermansia muciniphila</i> DSM 22959	702
354. Evaluation of the entomopathogenic fungus, <i>Beauveria bassiana</i> as a biocontrol agent against the chestnut moth, <i>Cydia splendana</i> in the Minho region	704
408. Using yeast as a model, synthetic cannabinoids consumption might be a risk factor on the onset of Parkinson's Disease	705
415. New <i>in vitro</i> models for research and drug discovery against hepatotropic pathogens	706

5.4 Digital Data, Big Data & AI in Microbiotech **708**

256. Counter-Propagation-Artificial Neural Network and FT-MIR spectroscopy: the proper strategy for detailed antimicrobial activity analysis	708
--	-----

Microbiotec Sponsors **710**

Organizing Committee

Chairs:

Cecília Roque, FCT-NOVA

Miguel Viveiros, IHMT-NOVA

Universidade NOVA Lisboa

Ana Coroadinha, ITQB/iBET-NOVA

Cecilia Roque, FCT-NOVA (Chair)

Celso Cunha, IHMT-NOVA

Filomena Freitas, FCT-NOVA

Isabel Couto, IHMT-NOVA

Maria Jesus Chasqueira, NMS-NOVA

Miguel Viveiros, IHMT-NOVA (Chair)

Raquel Sá-Leão, ITQB-NOVA

Rui Oliveira, FCT-NOVA

Sandra Tenreiro, NMS-NOVA

Scientific Committee

Ana Azevedo (IST, Universidade de Lisboa)

Ana Cristina Dias Cabral (Universidade da Beira Interior)

Ana Teresa Caldeira (HERCULES, Universidade de Évora)

Ângela Novais (FF/Universidade do Porto)

Artur Alves (CESAM, Universidade de Aveiro)

Carlos Martins (FMV, Universidade de Lisboa)

Cecília Leão (ECS, Universidade do Minho)

Célia Manaia (ESB, Universidade Católica Portuguesa)

Célia Silva (Universidade dos Açores)

Eugénio Ferreira (CEB, Universidade do Minho)

Gabriel Monteiro (IST, Universidade de Lisboa)

Helena Santos (ITQB, Universidade Nova de Lisboa)

Hermínia Lencastre (ITQB, Universidade Nova de Lisboa)

Isabel Sá Correia (IST, Universidade de Lisboa)
Jaime Mota (UCIBIO/FCT/Universidade Nova de Lisboa)
João Mano (CICECO, Universidade de Aveiro)
João Queiroz (Universidade da Beira Interior)
Joaquim Cabral (IST, Universidade de Lisboa)
Jorge Pedrosa (ICVS, Universidade do Minho)
Jorge Pereira (DEQ, Universidade de Coimbra)
Jorge Rocha (DEQ, Universidade de Coimbra)
José Teixeira (CEB, Universidade do Minho)
Lucília Domingues (CEB, Universidade do Minho)
Luísa Peixe (FF, Universidade do Porto)
Madalena Pimentel (FF, Universidade de Lisboa)
Manuela Pintado (ESB, Universidade Católica Portuguesa)
Mara Freire (CICECO, Universidade de Aveiro)
Margarida Casal (CBMA, Universidade do Minho)
Maria Ascensão Reis (UCIBIO/FCT/Universidade Nova de Lisboa)
Maria João Amorim (Instituto Gulbenkian de Ciência)
Maria José Saavedra (CITAB, Universidade de Trás-os-Montes e Alto Douro)
Maria Leonor Faleiro (Universidade do Algarve)
Mariana Pinho (ITQB/Universidade Nova de Lisboa)
Miguel Teixeira (IST, Universidade de Lisboa)
Mónica Vieira Cunha (FC, Universidade de Lisboa)
Nelson Lima (CEB, Universidade do Minho)
Nelson Simões (CBA, Universidade dos Açores)
Nuno Cerca (CEB, Universidade do Minho)
Paula Castro (ESB, Universidade Católica Portuguesa)
Paula Morais (CEMMPRE, Universidade de Coimbra)
Pedro Barros Fernandes (Universidade Lusófona)
Raquel Aires Barros (IST, Universidade de Lisboa)
Teresa Crespo (ITQB/iBET, Universidade Nova de Lisboa)
Teresa Gonçalves (CNC/FM, Universidade de Coimbra)
Vítor Vasconcelos (CIIMAR, Universidade do Porto)

PCO Secretariat

Organideia – Organização Profissional de Congressos, Lda.

Av. da Guarda Inglesa, 27

3040-193 Coimbra, PORTUGAL

Phone: +351 239 801 009

e-Mail: microbiotec21@organideia.pt

105. Aerobic granular sludge, a feasible technology for the treatment and recirculation of trout aquaculture water streams

Ana T. Couto¹, Catarina L. Amorim¹ and Paula M.L. Castro¹

¹*Universidade Católica Portuguesa, CBQF - Centro de Biotecnologia e Química Fina Laboratório Associado, Escola Superior de Biotecnologia, Porto, Portugal*

Aquaculture is one of the fastest growing food production sectors. However, land-based aquaculture faces water scarcity and space restrictions which hamper its development. Aquaculture recirculation systems (RAS) help tackle these problems, although in most systems the recycling of water increases nutrient and organic matter concentrations, which often translates to fish mortality.

This work aimed to evaluate the feasibility of an aerobic granular sludge (AGS) system as a novel alternative for the treatment of aquaculture streams, characterized by low nutrient and carbon concentrations. The efficiency for nutrients removal aiming at water recirculation was assessed as well as the AGS microbiome composition dynamics over operation.

The AGS system was inoculated with mature granules from a full-scale WWTP and fed with a synthetic medium mimicking a trout farm aquaculture recycling water. High ammonium removal efficiencies of 87-100% and ammonium removal rate of approximately 14.5 mg NH₄⁺-N/(L·d) were achieved. The AGS microbiome, mainly composed by members of the Proteobacteria phylum, had a core microbiome composed of different functional groups (e.g. denitrifying bacteria, nitrogen fixing bacteria, ammonium oxidizing bacteria, phosphorus accumulating bacteria and extracellular polymeric substance (EPS) producing bacteria).

The AGS system was able to produce effluents of high-quality, with ammonium content below the toxic levels for fish, and thus suitable for recirculating in the aquaculture.

Acknowledgments: The authors thank the EU and the FCT (Fundação para a Ciência e Tecnologia; Water JPI/0003/2016) for funding, in the frame of the collaborative international Consortium AquaVal financed under the ERA-NET WaterWorks2015 Cofunded Call. This ERA-NET is an integral part of the 2016 Joint Activities developed by the Water Challenges for a Changing World Joint Programme Initiative (Water JPI). Authors also thank the CBQF collaboration under the FCT project UIDB/50016/2020. A.T. Couto thanks the research grant from FCT (SFRH/BD/139924/2018) and POCH, supported by the European Social Fund and MCTES national funds.